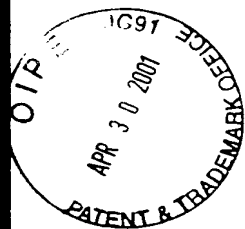


--The Sequence Listing written in file -140.APP, 582,722 bytes, created on April 26, 2001 on two identical copies of compact discs for Application No: 09/357,737, Sette *et al.*, INDUCING CELLULAR RESPONSES TO HEPATITIS C VIRUS USING PEPTIDE AND NUCLEIC ACID COMPOSITIONS, is hereby incorporated-by-reference.--

Please replace the paragraph beginning at page 76, line 5, with the following:

--In summary, on the basis of the data presented in the above examples, 26 CTL candidate peptide epitopes derived from conserved regions of the HCV virus have been identified (Table XXXVIa). These include twelve HLA-A2 supermotif-bearing epitopes, eight HLA-A3 supermotif-bearing epitopes, and one HLA-B7 supermotif-bearing epitope, each capable of binding to multiple A2-, A3-, or B7-supertype molecules, and immunogenic in HLA transgenic mice or antigenic for human PBL (with the exception of peptide 29.0035/1260.04). Additional epitopes not evaluated for immunogenicity are also included. They are an additional B7-supermotif-bearing epitope and two HLA-A1 and one HLA-A24 high-affinity binding peptides. A known HLA-A31 restricted epitope (VGIYLLPNR; SEQ ID NO:3587), which also binds HLA-A33, is also set out in Table XXXVIa and is useful in combination with other Class I or Class II epitopes.--

Please replace the paragraph beginning at page 99, line 1 (TABLE III), with the following:



--TABLE III

POSITION

MOTIFS	POSITION								
	1° anchor 1	2	3	4	5	1° anchor 6	7	8	9
DR4 preferred	FMYLIVW	M	T		I	VSTCPALIM	MH		MH
deleterious				W			R		WDE
DR1 preferred	MFLIVWY			PAMQ		VMATSPLIC	M		AVM
deleterious		C	CH	FD	CWD		GDE	D	
DR7 preferred	MFLIVWY	M	W	A		IVMSACTPL	M		IV (SEQ ID NO:3682)
deleterious		C		G			GRD	N	G (SEQ ID NO:3683)
DR Supermotif	MFLIVWY					VMSTACPLI			
DR3 MOTIFS	1° anchor 1	2	3		1° anchor 4	5	1° anchor 6		
motif 'a' preferred	LIVMFY				D				
motif 'b' preferred	LIVMFAY				DNQUEST		KRIH		

Italicized residues indicate less preferred or "tolerated" residues.--